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Key indicators

Single-crystal X-ray study
 $T = 283\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$
 R factor = 0.031
 wR factor = 0.094
Data-to-parameter ratio = 15.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Poly[[$(1,10\text{-phenanthroline-}\kappa^2\text{N,N'})\text{cadmium(II)}\text{-}\mu_3\text{-pyridine-3,4\text{-dicarboxylato-}\kappa^5\text{N:O,O':O'',O'''}]$]

In the title compound, $[\text{Cd}(\text{C}_7\text{H}_3\text{NO}_4)(\text{C}_{12}\text{H}_8\text{N}_2)]_n$, the Cd^{II} atom has a distorted pentagonal bipyramidal coordination geometry, defined by three N atoms and four carboxyl O atoms from chelating 1,10-phenanthroline and pyridine-3,4-dicarboxylate ligands. The structure exhibits a novel two-dimensional wave-like layer parallel to the (110) plane, with (6,3)-topology, which further forms a three-dimensional supramolecular network *via* π - π stacking.

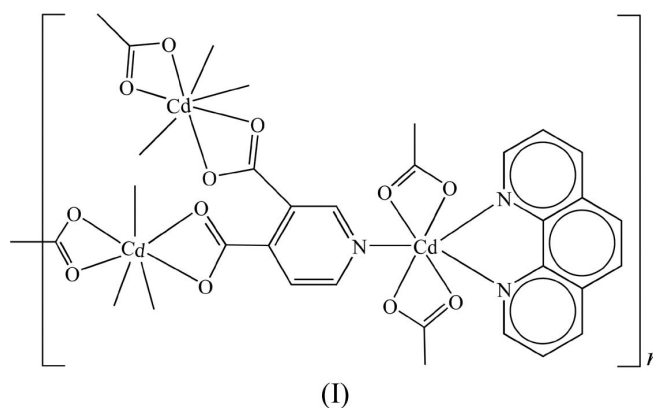
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Comment

In the last few years, the rational design and synthesis of metal-organic polymers has been of interest in the field of supramolecular chemistry and crystal engineering. The 3,4-pyridinedicarboxylate anion (PDB) has been investigated in the construction of metal-organic frameworks. Some structures of metal-organic polymers containing PDB have been reported previously (Wang, Qin, Wang, Li, Hao *et al.*, 2004; Wang, Qin, Wang, Li, Hu & Xu, 2004; Tong *et al.*, 2005; Zhang *et al.*, 2005; Chen *et al.*, 2003; Xia *et al.*, 2004; Qin *et al.*, 2005). Of these, two complexes, namely PDB-(2,2'-bipyridine)zinc(II) and PDB-(2,2'-bipyridine)cadmium(II), display a layered structure (Qin *et al.*, 2005). We present here the crystal structure of the title compound, (I), which is a new example of a compound with a layered polymeric structure.



As shown in Fig. 1, the asymmetric unit of (I) contains one Cd^{II} atom, one PDB and one 1,10-phenanthroline (phen) ligand. The Cd^{II} atom is coordinated by four O atoms from two PDB ligands and three N atoms from PDB and phen ligands, showing a distorted pentagonal bipyramidal coordination geometry. The distortion is considerable because of the small bite angles [$\text{O}-\text{Cd}-\text{O}$ angles = $52.38(8)$ and $53.23(8)^\circ$] when PDB ligands coordinate with metal atoms. This is very different from the distorted square-pyramidal

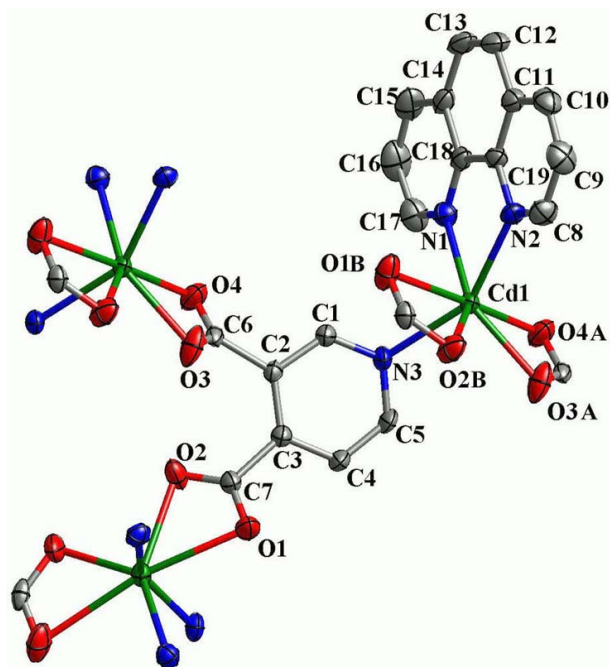


Figure 1
The molecular structure of (I), showing the atom-numbering scheme and with 50% probability displacement ellipsoids. H atoms have been omitted. [Symmetry codes: (A) $1 + x, y, z$; (B) $-x, \frac{1}{2} + y, \frac{1}{2} - z$.]

geometry found in PDB-(2,2'-bipyridine)cadmium(II) (Qin *et al.*, 2005). The Cd—O bond distances range from 2.262 (3) to 2.645 (3) Å, while the Cd—N bond distances range from 2.346 (3) to 2.388 (3) Å (Table 1).

Adjacent Cd^{II} atoms are linked by the PDB ligands to form a two-dimensional wave-like layer parallel to the (110) plane. Each PDB ligand serves as a three-connector, connecting three Cd^{II} atoms, while the Cd^{II} atom acts as a tri-connected node. The spontaneous association of these subunits results in a new kind of two-dimensional hexagonal (6,3) sheet, the three-connecting nodes of which are provided by both PDB²⁻ and Cd²⁺ (Fig. 2). The phen ligand chelates to the Cd^{II} atom in the direction nearly perpendicular to the two-dimensional layer. This orientation plays an important role in the packing. A three-dimensional network is formed *via* π - π stacking interactions between phen ligands, the face-to-face distances between neighbouring parallel phen planes being 3.436 (3) and 3.571 (3) Å (Fig. 3).

Experimental

Compound (I) was prepared by a hydrothermal method. A mixture of Cd(NO₃)₂·6H₂O (0.5 mmol), H₂PDB (0.6 mmol) and water (10 ml) was stirred for 20 min in air. The mixture was then transferred to a 23 ml Teflon reactor and kept at 433 K for 72 h under autogenous pressure. Single crystals of (I) suitable for X-ray analysis were obtained from the reaction mixture.

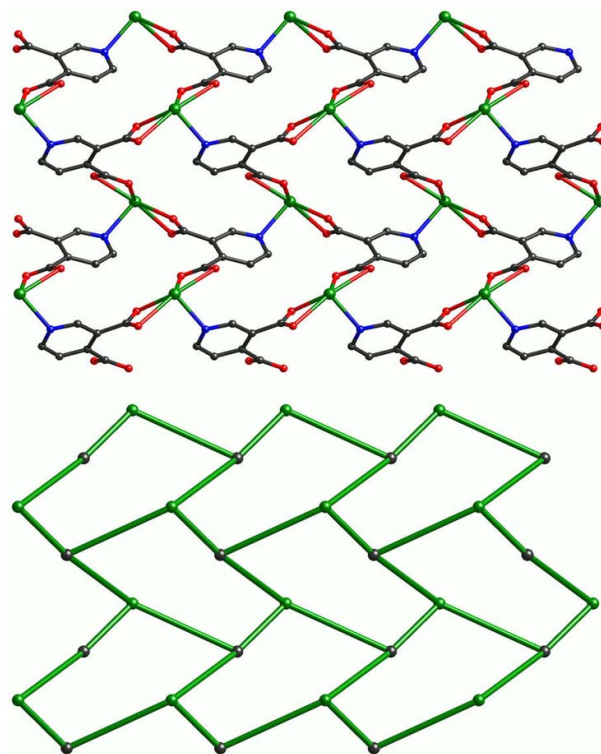


Figure 2
A perspective view of the two-dimensional layer, with phen ligands and H atoms omitted (top) and the (6,3) net (bottom).

Crystal data

[Cd(C₇H₃NO₄)(C₁₂H₈N₂)]
M_r = 457.71
 Monoclinic, *P*2₁/*c*
a = 7.5743 (15) Å
b = 9.4265 (19) Å
c = 23.070 (5) Å
 β = 99.17 (3)°
V = 1626.1 (6) Å³
Z = 4

D_x = 1.870 Mg m⁻³
 Mo *K* α radiation
 Cell parameters from 15350 reflections
 θ = 3.0–27.5°
 μ = 1.38 mm⁻¹
T = 283 (2) K
 Block, colourless
 0.22 × 0.20 × 0.16 mm

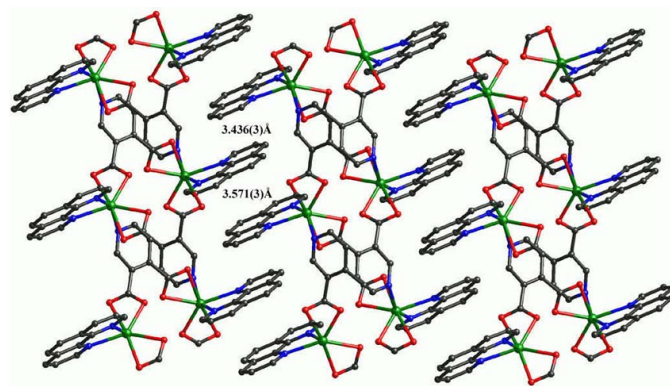


Figure 3
A packing model of the three-dimensional supramolecular network. H atoms have been omitted.

Data collection

Rigaku R-AXIS RAPID
diffractometer
 ω scans
Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)
 $T_{\min} = 0.752$, $T_{\max} = 0.810$
15350 measured reflections

3700 independent reflections
3190 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.034$
 $\theta_{\text{max}} = 27.5^\circ$
 $h = -9 \rightarrow 9$
 $k = -12 \rightarrow 12$
 $l = -27 \rightarrow 29$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.031$
 $wR(F^2) = 0.094$
 $S = 1.01$
3700 reflections
244 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.065P)^2 + 0.473P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.48 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.65 \text{ e } \text{\AA}^{-3}$

Table 1

Selected bond lengths (\AA).

| | | | |
|---------------------|-----------|----------------------|-----------|
| Cd1—N1 | 2.360 (3) | Cd1—O2 ⁱ | 2.406 (2) |
| Cd1—N2 | 2.346 (3) | Cd1—O3 ⁱⁱ | 2.262 (3) |
| Cd1—N3 | 2.388 (3) | Cd1—O4 ⁱⁱ | 2.645 (3) |
| Cd1—O1 ⁱ | 2.494 (3) | | |

Symmetry codes: (i) $-x, y + \frac{1}{2}, -z + \frac{1}{2}$; (ii) $x + 1, y, z$.

The H atoms were placed in calculated positions, with C—H = 0.93 \AA , and treated as riding atoms in the final cycles of refinement, with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *PROCESS-AUTO* (Rigaku, 1998); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSK, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *CrystalStructure*; software used to prepare material for publication: *SHELXL97*.

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